

* * * * * STN Columbus * * * * *

FILE 'HOME' ENTERED AT 12:01:09 ON 25 JUL 2006

=> file medline caplus biosis

COST IN U.S. DOLLARS

SINCE FILE

TOTAL

ENTRY

SESSION

FULL ESTIMATED COST

0.21

0.21

FILE 'MEDLINE' ENTERED AT 12:01:26 ON 25 JUL 2006

FILE 'CAPLUS' ENTERED AT 12:01:26 ON 25 JUL 2006

USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.

PLEASE SEE "HELP USAGETERMS" FOR DETAILS.

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FILE 'BIOSIS' ENTERED AT 12:01:26 ON 25 JUL 2006

Copyright (c) 2006 The Thomson Corporation

=> s (array or microarray) (p) ("small RNA" or smRNA or microRNA or snoRNA or sRNA or ncRNA or nmR
L1 414 (ARRAY OR MICROARRAY) (P) ("SMALL RNA" OR SMRNA OR MICRORNA OR
SNORNA OR SRNA OR NCRNA OR NMRNA OR SNMRNA OR FRNA OR SNRNA)

=> dup remove l1

PROCESSING COMPLETED FOR L1

L2 227 DUP REMOVE L1 (187 DUPLICATES REMOVED)

=> s (array or microarray) (8a) ("small RNA" or smRNA or microRNA or snoRNA or sRNA or ncRNA or nm
L3 176 (ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA OR
SNORNA OR SRNA OR NCRNA OR NMRNA OR SNMRNA OR FRNA OR SNRNA)

=> dup remove l3

PROCESSING COMPLETED FOR L3

L4 105 DUP REMOVE L3 (71 DUPLICATES REMOVED)

=> s (array or microarray) (8a) ("small RNA" or smRNA or microRNA or snoRNA or sRNA or ncRNA or nm
L5 8 (ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA OR
SNORNA OR SRNA OR NCRNA OR NMRNA OR SNMRNA OR FRNA OR SNRNA)
(8A) (PERFECT OR MISMATCH)

=> dup remove l5

PROCESSING COMPLETED FOR L5

L6 3 DUP REMOVE L5 (5 DUPLICATES REMOVED)

=> d ti 1-3

L6 ANSWER 1 OF 3 MEDLINE on STN DUPLICATE 1
TI Concerted evolution of the tandem array encoding primate U2 snRNA (the
RNU2 locus) is accompanied by dramatic remodeling of the junctions with
flanking chromosomal sequences.

L6 ANSWER 2 OF 3 MEDLINE on STN DUPLICATE 2
TI Concerted evolution of the tandemly repeated genes encoding primate U2
small nuclear RNA (the RNU2 locus) does not prevent rapid diversification
of the (CT)n.(GA)n microsatellite embedded within the U2 repeat unit.

L6 ANSWER 3 OF 3 MEDLINE on STN DUPLICATE 3
TI Human genes for U2 small nuclear RNA are tandemly repeated.

=> s ((array or microarray) (8a) ("small RNA" or smRNA or microRNA or snoRNA or sRNA or ncRNA or n
L7 0 ((ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA
OR SNORNA OR SRNA OR NCRNA OR NMRNA OR SNMRNA OR FRNA OR SNRNA))
AND ((MISMATCH OR PERFECT) (3A) PROBE)

=> s ((array or microarray) AND ("small RNA" or smRNA or microRNA or snoRNA or sRNA or ncRNA or nm
L8 0 ((ARRAY OR MICROARRAY) AND ("SMALL RNA" OR SMRNA OR MICRORNA OR
SNORNA OR SRNA OR NCRNA OR NMRNA OR SNMRNA OR FRNA OR SNRNA))
AND ((MISMATCH OR PERFECT) (3A) PROBE)

=> d his

(FILE 'HOME' ENTERED AT 12:01:09 ON 25 JUL 2006)

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FILE 'MEDLINE, CAPLUS, BIOSIS' ENTERED AT 12:01:26 ON 25 JUL 2006
L1      414 S (ARRAY OR MICROARRAY) (P) ("SMALL RNA" OR SMRNA OR MICRORNA O
L2      227 DUP REMOVE L1 (187 DUPLICATES REMOVED)
L3      176 S (ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA
L4      105 DUP REMOVE L3 (71 DUPLICATES REMOVED)
L5      8 S (ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA
L6      3 DUP REMOVE L5 (5 DUPLICATES REMOVED)
L7      0 S ((ARRAY OR MICROARRAY) (8A) ("SMALL RNA" OR SMRNA OR MICRORNA
L8      0 S ((ARRAY OR MICROARRAY) AND ("SMALL RNA" OR SMRNA OR MICRORNA

```

```

=> s l2 and ((perfect or mismatch) (5a) probe)
L9      0 L2 AND ((PERFECT OR MISMATCH) (5A) PROBE)

```

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=> s l2 and ("perfect match probe" or "mismatch probe")
L10     0 L2 AND ("PERFECT MATCH PROBE" OR "MISMATCH PROBE")

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=> logoff

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* * * * * STN Columbus * * * * *

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FILE 'HOME' ENTERED AT 11:25:32 ON 26 JUL 2006

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=> index all

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FILE 'ENCOMPLIT2' ACCESS NOT AUTHORIZED

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FILE 'ENCOMPAT2' ACCESS NOT AUTHORIZED

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COST IN U.S. DOLLARS	SINCE FILE ENTRY	TOTAL SESSION
FULL ESTIMATED COST	0.21	0.21

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INDEX 'ABI-INFORM, ADISCTI, AEROSPACE, AGRICOLA, ALUMINIUM, ANABSTR, ANTE,
      APOLLIT, AQUALINE, AQUASCI, AQUIRE, BABS, BIBLIODATA, BIOENG, BIOSIS,
      BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAOLD, CAPLUS, CASREACT, CBNB,
      CEABA-VTB, CERAB, CHEMINFORMRX, CHEMSAFE, ...'

```

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ENTERED AT 11:25:49 ON 26 JUL 2006

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135 FILES IN THE FILE LIST IN STNINDEX

```

```

Enter SET DETAIL ON to see search term postings or to view
search error messages that display as 0* with SET DETAIL OFF.

```

```

=> s ((array or microarray) (p) ("perfect match probe" or "PM probe") and (microRNA or sRNA or sno
MISMATCHED QUOTE 'OR "SMALL'
Quotation marks (or apostrophes) must be used in pairs,
one before and one after the expression you are setting
off or masking.

```

```

=> s ((array or microarray) (p) ("perfect match probe" or "PM probe") and (microRNA or sRNA or sno
UNMATCHED LEFT PARENTHESIS '((ARRAY'
The number of right parentheses in a query must be equal to the
number of left parentheses.

```

```

=> s (array or microarray) (p) ("perfect match probe" or "PM probe") and (microRNA or sRNA or snoR
0* FILE ALUMINIUM
0* FILE ANTE
0* FILE APOLLIT
<-----User Break----->

```

```

=> s (array or microarray) (8a) ("perfect match probe" or "PM probe" or affymetrix) and (microRNA
1 FILE AGRICOLA
4 FILE BIOSIS
3 FILE BIOTECHABS
3 FILE BIOTECHDS
1 FILE CABA
20 FILES SEARCHED...
22 FILE CAPLUS
29 FILES SEARCHED...
1 FILE COMPENDEX
42 FILES SEARCHED...
1 FILE DISSABS
2 FILE EMBASE
5 FILE EPFULL
4 FILE ESBIODBASE
57 FILES SEARCHED...
3 FILE GBFULL

```

67 FILES SEARCHED...
1 FILE IFIPAT
83 FILES SEARCHED...
1 FILE LIFESCI
3 FILE MEDLINE
1 FILE PAPERCHEM2
1 FILE PASCAL

99 FILES SEARCHED...
126 FILE PCTFULL

111 FILES SEARCHED...

<-----User Break----->

=> file caplus

COST IN U.S. DOLLARS

SINCE FILE

TOTAL

ENTRY

SESSION

FULL ESTIMATED COST

5.49

5.70

FILE 'CAPLUS' ENTERED AT 11:31:06 ON 26 JUL 2006

USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.

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FILE COVERS 1907 - 26 Jul 2006 VOL 145 ISS 5

FILE LAST UPDATED: 25 Jul 2006 (20060725/ED)

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<http://www.cas.org/infopolicy.html>

=> s (array or microarray) (8a) ("perfect match probe" or "PM probe" or affymetrix) and (microRNA

117136 ARRAY

49947 ARRAYS

146899 ARRAY

(ARRAY OR ARRAYS)

39212 MICROARRAY

10114 MICROARRAYS

41618 MICROARRAY

(MICROARRAY OR MICROARRAYS)

35841 "PERFECT"

30 "PERFECTS"

35868 "PERFECT"

("PERFECT" OR "PERFECTS")

26068 "MATCH"

9949 "MATCHES"

34743 "MATCH"

("MATCH" OR "MATCHES")

226574 "PROBE"

113583 "PROBES"

299509 "PROBE"

("PROBE" OR "PROBES")

19 "PERFECT MATCH PROBE"

("PERFECT" (W) "MATCH" (W) "PROBE")

39869 "PM"

3610 "PMS"

43134 "PM"

("PM" OR "PMS")

226574 "PROBE"

113583 "PROBES"

299509 "PROBE"

("PROBE" OR "PROBES")

11 "PM PROBE"

("PM" (W) "PROBE")

2709 AFFYMETRIX

2150 (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROBE"
 OR AFFYMETRIX)
 1562 MICRORNA
 851 MICRORNAS
 1707 MICRORNA
 (MICRORNA OR MICRORNAS)
 833 SRNA
 90 SRNAS
 851 SRNA
 (SRNA OR SRNAS)
 493 SNORNA
 373 SNORNAS
 587 SNORNA
 (SNORNA OR SNORNAS)
 2036 SNRNA
 711 SNRNAS
 2277 SNRNA
 (SNRNA OR SNRNAS)
 86 NCRNA
 168 NCRNAS
 202 NCRNA
 (NCRNA OR NCRNAS)
 19 SMRNA
 7 SMRNAS
 21 SMRNA
 (SMRNA OR SMRNAS)
 7 NMRNA
 1 NMRNAS
 8 NMRNA
 (NMRNA OR NMRNAS)
 58 FRNA
 6 FRNAS
 61 FRNA
 (FRNA OR FRNAS)
 1225202 "SMALL"
 45 "SMALLS"
 1225241 "SMALL"
 ("SMALL" OR "SMALLS")
 311037 "RNA"
 25432 "RNAS"
 315711 "RNA"
 ("RNA" OR "RNAS")
 2052 "SMALL RNA"
 ("SMALL" (W) "RNA")
 L1 22 (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROBE"
 OR AFFYMETRIX) AND (MICRORNA OR SRNA OR SNORNA OR SNRNA OR NCRNA
 OR SMRNA OR NMRNA OR FRNA OR "SMALL RNA")

=> d ti 1-10

L1 ANSWER 1 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Optimization and validation of small quantity RNA profiling for
 identifying TNF responses in cultured human vascular endothelial cells
 L1 ANSWER 2 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Analysis of Affymetrix GeneChip data using amplified RNA
 L1 ANSWER 3 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI DICER-LIKE 4 functions in trans-acting small interfering RNA biogenesis
 and vegetative phase change in Arabidopsis thaliana
 L1 ANSWER 4 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Gene expression profiles in the diagnosis and treatment of Alzheimer's
 disease
 L1 ANSWER 5 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Gene family analysis of the Arabidopsis pollen transcriptome reveals
 biological implications for cell growth, division control, and gene
 expression regulation
 L1 ANSWER 6 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Gene expression profiles and biomarkers for the detection of
 hyperlipidemia and other disease-related gene transcripts in blood
 L1 ANSWER 7 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN

TI Gene expression profiles and biomarkers for the detection of hyperlipidemia and other disease-related gene transcripts in blood

L1 ANSWER 8 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN

TI Gene expression profiles and biomarkers for the detection of hyperlipidemia and other disease-related gene transcripts in blood

L1 ANSWER 9 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN

TI Gene expression profiles and biomarkers for the detection of Chagas disease and other disease-related gene transcripts in blood

L1 ANSWER 10 OF 22 CAPLUS COPYRIGHT 2006 ACS on STN

TI Gene expression profiles and biomarkers for the detection of Chagas disease and other disease-related gene transcripts in blood

```
=> s (array or microarray) (8a) ("perfect match probe" or "PM probe") and (microRNA or sRNA or sno
117136 ARRAY
49947 ARRAYS
146899 ARRAY
      (ARRAY OR ARRAYS)
39212 MICROARRAY
10114 MICROARRAYS
41618 MICROARRAY
      (MICROARRAY OR MICROARRAYS)
35841 "PERFECT"
30 "PERFECTS"
35868 "PERFECT"
      ("PERFECT" OR "PERFECTS")
26068 "MATCH"
9949 "MATCHES"
34743 "MATCH"
      ("MATCH" OR "MATCHES")
226574 "PROBE"
113583 "PROBES"
299509 "PROBE"
      ("PROBE" OR "PROBES")
19 "PERFECT MATCH PROBE"
      ("PERFECT" (W) "MATCH" (W) "PROBE")
39869 "PM"
3610 "PMS"
43134 "PM"
      ("PM" OR "PMS")
226574 "PROBE"
113583 "PROBES"
299509 "PROBE"
      ("PROBE" OR "PROBES")
11 "PM PROBE"
      ("PM" (W) "PROBE")
6 (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROBE")
1562 MICRORNA
851 MICRORNAS
1707 MICRORNA
      (MICRORNA OR MICRORNAS)
833 SRNA
90 SRNAS
851 SRNA
      (SRNA OR SRNAS)
493 SNORNA
373 SNORNAS
587 SNORNA
      (SNORNA OR SNORNAS)
2036 SNRNA
711 SNRNAS
2277 SNRNA
      (SNRNA OR SNRNAS)
86 NCRNA
168 NCRNAS
202 NCRNA
      (NCRNA OR NCRNAS)
19 SMRNA
7 SMRNAS
21 SMRNA
      (SMRNA OR SMRNAS)
7 NMRNA \
```

```

1 NMRNAS
8 NMRNA
  (NMRNA OR NMRNAS)
58 FRNA
6 FRNAS
61 FRNA
  (FRNA OR FRNAS)
1225202 "SMALL"
45 "SMALLS"
1225241 "SMALL"
  ("SMALL" OR "SMALLS")
311037 "RNA"
25432 "RNAS"
315711 "RNA"
  ("RNA" OR "RNAS")
2052 "SMALL RNA"
  ("SMALL" (W) "RNA")
L2 0 (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROBE")
    AND (MICRORNA OR SRNA OR SNORNA OR SNRNA OR NCRNA OR SMRNA OR
    NMRNA OR FRNA OR "SMALL RNA")

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=> d his

(FILE 'HOME' ENTERED AT 11:25:32 ON 26 JUL 2006)

INDEX 'ABI-INFORM, ADISCTI, AEROSPACE, AGRICOLA, ALUMINIUM, ANABSTR,
ANTE, APOLLIT, AQUALINE, AQUASCI, AQUIRE, BABS, BIBLIODATA, BIOENG,
BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAOLD, CAPLUS, CASREACT,
CBNB, CEABA-VTB, CERAB, CHEMINFORMRX, CHEMSAFE, ...' ENTERED AT 11:25:49
ON 26 JUL 2006

SEA (ARRAY OR MICROARRAY) (P) ("PERFECT MATCH PROBE" OR "PM PRO

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-----
0* FILE ALUMINIUM
0* FILE ANTE
0* FILE APOLLIT
0* FILE AQUALINE

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SEA (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PR

```

-----
1 FILE AGRICOLA
4 FILE BIOSIS
3 FILE BIOTECHABS
3 FILE BIOTECHDS
1 FILE CABA
22 FILE CAPLUS
1 FILE COMPENDEX
1 FILE DISSABS
2 FILE EMBASE
5 FILE EPFULL
4 FILE ESBIODBASE
3 FILE GBFULL
1 FILE IFIPAT
1 FILE LIFESCI
3 FILE MEDLINE
1 FILE PAPERCHEM2
1 FILE PASCAL
126 FILE PCTFULL
3 FILE SCISEARCH

```

FILE 'CAPLUS' ENTERED AT 11:31:06 ON 26 JUL 2006

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L1 22 S (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROB
L2 0 S (ARRAY OR MICROARRAY) (8A) ("PERFECT MATCH PROBE" OR "PM PROB

```

=> dup remove l1

PROCESSING COMPLETED FOR L1

L3 13 DUP REMOVE L1 (9 DUPLICATES REMOVED)

=> d ti 1-13

```

L3 ANSWER 1 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
TI Optimization and validation of small quantity RNA profiling for
identifying TNF responses in cultured human vascular endothelial cells

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L3 ANSWER 2 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
TI Gene expression profiles in the diagnosis and treatment of Alzheimer's
disease

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L3 ANSWER 3 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Biomarkers of cyclin-dependent kinase modulation in cancer therapy

L3 ANSWER 4 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI DICER-LIKE 4 functions in trans-acting small interfering RNA biogenesis and vegetative phase change in Arabidopsis thaliana

L3 ANSWER 5 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Gene family analysis of the Arabidopsis pollen transcriptome reveals biological implications for cell growth, division control, and gene expression regulation

L3 ANSWER 6 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Analysis of Affymetrix GeneChip data using amplified RNA

L3 ANSWER 7 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 1
 TI Gene expression profiles and biomarkers for the detection of depression-related and other disease-related gene transcripts in blood

L3 ANSWER 8 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 2
 TI Gene expression profiles and biomarkers for the detection of hyperlipidemia and other disease-related gene transcripts in blood

L3 ANSWER 9 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 3
 TI Gene expression profiles and biomarkers for the detection of Chagas disease and other disease-related gene transcripts in blood

L3 ANSWER 10 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 4
 TI Gene expression profiles and biomarkers for the detection of lung disease-related and other disease-related gene transcripts in blood

L3 ANSWER 11 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Gene expression profiling in predicting drug sensitivity in patients afflicted with hypertension

L3 ANSWER 12 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI Signature genes expressed the lung during asthma or allergies and their use in predicting, diagnosing and treating asthma or allergies

L3 ANSWER 13 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 TI GeneChip analysis of hippocampal gene expression profiles of short- and long-attack-latency mice: Technical and biological implications

=> d bib kwic 1

L3 ANSWER 1 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
 AN 2006:178532 CAPLUS <<LOGINID::20060726>>
 DN 144:329540
 TI Optimization and validation of small quantity RNA profiling for identifying TNF responses in cultured human vascular endothelial cells
 AU Shou, Jianyong; Qian, Hui-Rong; Lin, Xi; Stewart, Trent; Onyia, Jude E.; Gelbert, Lawrence M.
 CS Integrative Biology, Lilly Research Laboratories, Indianapolis, IN, 46285, USA
 SO Journal of Pharmacological and Toxicological Methods (2006), 53(2), 152-159
 CODEN: JPTMEZ; ISSN: 1056-8719
 PB Elsevier B.V.
 DT Journal
 LA English

RE.CNT 23 THERE ARE 23 CITED REFERENCES AVAILABLE FOR THIS RECORD
 ALL CITATIONS AVAILABLE IN THE RE FORMAT

AB ***Affymetrix*** oligonucleotide ***microarrays*** are widely used in basic and applied research. The need for a significant amt. of starting RNA has limited its use in applications where the amt. of RNA is limiting, such as with Laser Captured Microdissection (LCM), small biopsies, or peripheral blood in rodent models. To overcome this limitation, various RNA amplification and labeling methods have been described, however, further optimization and validation of these methods are needed. Here we reported using the Arcturus technol. to optimize amplification and labeling of small amts. of RNA for ***Affymetrix*** ***microarray*** studies. We assessed the tech. feasibility and variation introduced by differences in starting RNA quantity and

differences in tech. performance by microarray hybridization. We demonstrated that the current approach is reliable to amplify as little as 40 ng total RNA, and it is suitable for Affymetrix studies yielding satisfactory quant. chip performance. We also showed that differences in labeling methods contribute more to variation than the differences in starting RNA quantity per se. As a model, we studied the well-documented TNF-induced inflammatory responses in cultured human vascular endothelial cells. We were able to recapitulate the TNF-induced responses using
small ***RNA*** sample profiling.

=> d kwic 13

L3 ANSWER 13 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
AB present study is to extend and validate the SAGE expression profile of hippocampus of LAL and SAL mice using GeneChips (***Affymetrix***, Santa Clara, CA; one ***array*** per mouse, n = 5 per mouse line). As was the case with SAGE, GeneChips detect only medium- to high-abundance. . . (gas5), in LAL mice. As gas5 does not encode a protein but several small nuclear RNAs, our data suggest that ***small*** ***RNAs*** may contribute to the mol. mechanisms underlying the extreme behavioral differences between LAL and SAL mice.

=> d bib 13

L3 ANSWER 13 OF 13 CAPLUS COPYRIGHT 2006 ACS on STN
AN 2003:963801 CAPLUS <<LOGINID::20060726>>
DN 140:123529
TI GeneChip analysis of hippocampal gene expression profiles of short- and long-attack-latency mice: Technical and biological implications
AU Feldker, Dorine E. M.; Datson, Nicole A.; Veenema, Alexa H.; Proutski, Vitali; Lathouwers, Desiree; de Kloet, E. Ron; Vreugdenhil, Erno
CS Division of Medical Pharmacology, Leiden/Amsterdam Center for Drug Research, Leiden University Medical Center, Leiden, Neth.
SO Journal of Neuroscience Research (2003), 74(5), 701-716
 CODEN: JNREDK; ISSN: 0360-4012
PB Wiley-Liss, Inc.
DT Journal
LA English
RE.CNT 32 THERE ARE 32 CITED REFERENCES AVAILABLE FOR THIS RECORD
 ALL CITATIONS AVAILABLE IN THE RE FORMAT

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